

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING

ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

10/031342A

Source:

INWIG

Date Processed by STIC:

6-7-05

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses

1. EFS-Bio (<http://www.uspto.gov/efb/efs/downloads/documents.htm>) , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window - Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/24/05



IFW16

RAW SEQUENCE LISTING

DATE: 06/07/2005

PATENT APPLICATION: US/10/031,342A

TIME: 10:29:35

Input Set : A:\2005-05-31 Sub SeqList.ST25.txt.txt

Output Set: N:\CRF4\06072005\J031342A.raw

3 <110> APPLICANT: M&E Biotech A/S
 5 <120> TITLE OF INVENTION: Method for down-regulating GDF-8 activity
 7 <130> FILE REFERENCE: AutoVacGDF-8 DK 1
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/031,342A
 C--> 9 <141> CURRENT FILING DATE: 2002-03-26
 E--> 9 <160> NUMBER OF SEQ ID NOS: 23 Found 24
 11 <170> SOFTWARE: PatentIn Ver. 2.1

ERRORED SEQUENCES

1414 <210> SEQ ID NO: 24
 1415 <211> LENGTH: 13
 1416 <212> TYPE: PRT
 1417 <213> ORGANISM: artificial sequence
 W--> 1419 <220> FEATURE:
 W--> 1419 <223> OTHER INFORMATION:
 E--> 1419 <400> SEQUENCE: 13-24
 1420 Ala Lys Phe Val Ala Ala Trp Thr Leu Lys Ala Ala Ala
 1421 1 5 10

Does Not Comply
 Corrected Diskette Needed
 (pg.1)

See error explanation on page 2.

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/031,342A

DATE: 06/07/2005
TIME: 10:29:36

Input Set : A:\2005-05-31 Sub SeqList.ST25.txt.txt

Output Set: N:\CRF4\06072005\J031342A.raw

Use of <220> Feature(NEW RULES):

Sequence(s) are missing the <220> Feature and associated headings.

Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence" or "Unknown". Please explain source of genetic material in <220> to <223> section (See "Federal Register," 6/01/98, Vol. 63, No. 104, pp.29631-32) (Sec.1.823 of new Rules)

ERROR EXPLANATION: ?

Seq#:24

VERIFICATION SUMMARY

DATE: 06/07/2005

PATENT APPLICATION: US/10/031,342A

TIME: 10:29:36

Input Set : A:\2005-05-31 Sub SeqList.ST25.txt.txt

Output Set: N:\CRF4\06072005\J031342A.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:1419 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:13, <213>
ORGANISM:artificial sequence
L:1419 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:13, <213>
ORGANISM:artificial sequence
L:1419 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:24 differs:13
L:1419 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:24,Line#:1419
L:9 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (23) Counted (24)